



SEQUENCE LISTING

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TECH CENTER 1600/2900

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<120> BIOENGINEERED VEHICLES FOR TARGETED NUCLEIC ACID
DELIVERY

<130> 23611-A USA

<140> As yet unassigned

<141> 2001-06-25

<150> 60/213,653

<151> 2000-06-23

<160> 53

<170> PatentIn Ver. 3.1

<210> 1

<211> 18

<212> PRT

<213> Homo sapiens

<400> 1

Ser Arg Ser Arg Tyr Tyr Arg Gln Arg Gln Arg Ser Arg Arg Arg Arg

1

5

10

15

Arg Arg

B2

<210> 2

<211> 26

<212> PRT

<213> Homo sapiens

<400> 2

Ala Lys Lys Ala Lys Ser Pro Lys Lys Ala Lys Ala Ala Lys Pro Lys

1

5

10

15

Lys Ala Pro Lys Ser Pro Ala Lys Ala Lys

20

25

<210> 3

<211> 10

<212> PRT

<213> Adenovirus

<400> 3

Ser Gly Pro Ser Asn Thr Pro Pro Glu Ile

1

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<210> 4

<211> 9

<212> PRT

<213> Human papillomavirus

<400> 4

Arg Ala His Tyr Asn Ile Val Thr Phe

1

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<211> 10

<212> PRT

<213> Human papillomavirus

<400> 5

Thr Asp Leu Tyr Cys Tyr Glu Gln Leu Asn

1 5 10

<210> 6

<211> 10

<212> PRT

<213> Human papillomavirus

<400> 6

Ala Glu Pro Asp Arg Ala His Tyr Asn Ile

1 5 10

<210> 7

<211> 19

<212> PRT

<213> Human papillomavirus

<400> 7

Lys Cys Asp Ser Thr Leu Arg Leu Cys Val Gln Ser Thr His Val Ile

1 5 10 15

Arg Thr Leu

<210> 8

<211> 10

<212> PRT

<213> Human papillomavirus

<400> 8

Gly Thr Leu Gly Ile Val Cys Pro Ile Cys

1 5 10

B2
Out

<210> 9
 <211> 10
 <212> PRT
 <213> Epstein-Barr Virus

<400> 9
 Asp Thr Pro Leu Ile Pro Leu Thr Ile Phe
 1 5 10

<210> 10
 <211> 15
 <212> PRT
 <213> Epstein-Barr Virus

<400> 10
 Pro Arg Ser Pro Thr Val Phe Tyr Asn Ile Pro Pro Met Pro Leu
 1 5 10 15

<210> 11
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 <212> PRT
 <213> Epstein-Barr Virus

<400> 11
 Phe Leu Arg Gly Arg Ala Tyr Gly Leu
 1 5

<210> 12
 <211> 15
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 <213> Epstein-Barr Virus

<400> 12

Arg Gly Ile Lys Glu His Val Ile Gln Asn Ala Phe Arg Lys Ala
 1 5 10 15

<210> 13

<211> 10

<212> PRT

<213> Epstein-Barr Virus

<400> 13

Glu Glu Asn Leu Leu Asp Phe Val Arg Phe
 1 5 10

<210> 14

<211> 9

<212> PRT

<213> Epstein-Barr Virus

<400> 14

Ile Val Thr Asp Phe Ser Val Ile Lys
 1 5

<210> 15

<211> 9

<212> PRT

<213> Homo sapiens

<400> 15

Leu Leu Gly Arg Asn Ser Pro Glu Val
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<210> 16

<211> 13

<212> PRT

<213> Murine sarcoma virus

<400> 16
 Lys Leu Val Val Val Gly Ala Arg Gly Val Gly Lys Ser
 1 5 10

<210> 17
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 <212> PRT
 <213> Homo sapiens

<400> 17
 Lys Leu Val Val Val Gly Ala Val Gly Val Gly Lys
 1 5 10

<210> 18
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 <212> PRT
 <213> Homo sapiens

<400> 18
 Asp Ile Leu Asp Thr Ala Gly Leu Glu Glu Tyr Ser Ala Met Arg Asp
 1 5 10 15

<210> 19
 <211> 8
 <212> PRT
 <213> Homo sapiens

<400> 19
 Gly Leu Glu Glu Tyr Ser Ala Met
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<210> 20
 <211> 10
 <212> PRT

<213> Homo sapiens

<400> 20

Glu Leu Val Ser Glu Phe Ser Arg Met Ala

1 5 10

<210> 21

<211> 15

<212> PRT

<213> Homo sapiens

<400> 21

His Leu Asp Met Leu Arg His Leu Tyr Gln Gly Cys Gln Val Val

1 5 10 15

<210> 22

<211> 15

<212> PRT

<213> Homo sapiens

<400> 22

Ser Arg Leu Leu Gly Ile Cys Leu Thr Ser Thr Val Gln Leu Val

1 5 10 15

<210> 23

<211> 9

<212> PRT

<213> Homo sapiens

<400> 23

Glu Ala Asp Pro Thr Gly His Ser Tyr

1 5

<210> 24

Be
Out

<211> 10

<212> PRT

<213> Homo sapiens

<400> 24

Leu Leu Asp Gly Thr Ala Thr Leu Arg Leu

1 5 10

<210> 25

<211> 9

<212> PRT

<213> Homo sapiens

<400> 25

Tyr Leu Glu Pro Gly Pro Val Thr Ala

1 5

<210> 26

<211> 9

<212> PRT

<213> Homo sapiens

<400> 26

Met Leu Leu Ala Val Leu Tyr Cys Leu

1 5

<210> 27

<211> 9

<212> PRT

<213> Homo sapiens

<400> 27

Tyr Met Asn Gly Thr Met Ser Gln Val

1 5

<210> 28

<211> 9

<212> PRT

<213> Homo sapiens

<400> 28

Tyr Met Asn Gly Thr Met Ser Glu Val

1

5

<210> 29

<211> 21

<212> PRT

<213> Homo sapiens

<400> 29

Ala Ala Gly Ile Gly Ile Leu Thr Val Ile Leu Gly Val Leu Leu Leu

1

5

10

15

Ile Gly Cys Trp Tyr

20

<210> 30

<211> 9

<212> PRT

<213> Simian virus 40

<400> 30

Thr Pro Pro Lys Lys Lys Arg Lys Val

1

5

<210> 31

<211> 14

<212> PRT

<213> Homo sapiens

<400> 31

Lys Lys Ser Ala Lys Lys Thr Pro Lys Lys Ala Lys Lys Pro

1

5

10

<210> 32

<211> 26

<212> PRT

<213> Homo sapiens

<400> 32

Ala Lys Lys Ala Lys Ser Pro Lys Lys Ala Lys Ala Ala Lys Pro Lys

1

5

10

15

Lys Ala Pro Lys Ser Pro Ala Lys Ala Lys

20

25

<210> 33

<211> 18

<212> PRT

<213> Homo sapiens

<400> 33

Ser Arg Ser Arg Tyr Tyr Arg Gln Arg Gln Arg Ser Arg Arg Arg Arg

1

5

10

15

Arg Arg

<210> 34

<211> 255

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Human/murine

chimeric single chain binding polypeptide (C6.5
sFv)

<400> 34

Gln Val Gln Leu Leu Gln Ser Gly Ala Glu Leu Lys Lys Pro Gly Glu
1 5 10 15

Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe Thr Ser Tyr
20 25 30

Trp Ile Ala Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Tyr Met
35 40 45

Gly Leu Ile Tyr Pro Gly Asp Ser Asp Thr Lys Tyr Ser Pro Ser Phe
50 55 60

Gln Gly Gln Val Thr Ile Ser Val Asp Lys Ser Val Ser Thr Ala Tyr
65 70 75 80

Leu Gln Trp Ser Ser Leu Lys Pro Ser Asp Ser Ala Val Tyr Phe Cys
85 90 95

Ala Arg His Asp Val Gly Tyr Cys Ser Ser Ser Asn Cys Ala Lys Trp
100 105 110

Pro Glu Tyr Phe Gln His Trp Gly Gln Gly Thr Leu Val Thr Val Ser
115 120 125

Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
130 135 140

Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Ala Ala Pro Gly Gln
145 150 155 160

Lys Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Asn Asn
165 170 175

Tyr Val Ser Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
180 185 190

B2
Cont

Ile Tyr Gly His Thr Asn Arg Pro Ala Gly Val Pro Asp Arg Phe Ser
 195 200 205

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Phe Arg
 210 215 220

Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ala Trp Asp Asp Ser Leu
 225 230 235 240

Ser Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly
 245 250 255

<210> 35

<211> 765

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human/murine
 chimeric single chain binding polypeptide (C6.5
 sFv)

<400> 35

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 tcctgtaagg gttctggata cagctttacc agctactgga tcgcctgggt gcgccagatg 120
 cccgggaaag gcctggagta catggggctc atctatcctg gtgactctga caccaaatac 180
 agcccgctct tccaaggcca ggtcaccatc tcagtcgaca agtcgcgtcag cactgcctac 240
 ttgcaatgga gcagtctgaa gccctcggac agcgccgtgt atttttgtgc gagacatgac 300
 gtgggatatt gcagtagttc caactgcgca aagtggcctg aatacttcca gcattggggc 360
 cagggcaccc tggtcaccgt ctctcaggt ggaggcgggt caggcggagg tggctctggc 420
 ggtggcggat cgcagtctgt gttgacgcag ccgcctcag tgtctgcggc ccaggacag 480
 aaggtcacca tctcctgctc tggaagcagc tccaacattg ggaataatta tgtatcctgg 540
 taccagcagc tcccaggaac agccccaaa ctctcatct atggtcacac caatcggccc 600
 gcaggggtcc ctgaccgatt ctctggctcc aagtctggca cctcagcctc cctggccatc 660
 agtgggttcc ggtccgagga tgaggctgat tattactgtg cagcatggga tgacagcctg 720
 agtggttggg tggttcggcg aggaccaag ctgaccgtcc taggt 765

<210> 36

<211> 269

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human/murine
chimeric single chain binding polypeptide (C6ML3-9
sFv')

<400> 36

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
1 5 10 15

Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe Thr Ser Tyr
20 25 30

Trp Ile Ala Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Tyr Met
35 40 45

Gly Leu Ile Tyr Pro Gly Asp Ser Asp Thr Lys Tyr Ser Pro Ser Phe
50 55 60

Gln Gly Gln Val Thr Ile Ser Val Asp Lys Ser Val Ser Thr Ala Tyr
65 70 75 80

Leu Gln Trp Ser Ser Leu Lys Pro Ser Asp Ser Ala Val Tyr Phe Cys
85 90 95

Ala Arg His Asp Val Gly Tyr Cys Ser Ser Ser Asn Cys Ala Lys Trp
100 105 110

Pro Glu Tyr Phe Gln His Trp Gly Gln Gly Thr Leu Val Thr Val Ser
115 120 125

Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
130 135 140

B2
Cont

Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Ala Ala Pro Gly Gln
 145 150 155 160

Lys Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Asn Asn
 165 170 175

Tyr Val Ser Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
 180 185 190

Ile Tyr Asp His Thr Asn Arg Pro Ala Gly Val Pro Asp Arg Phe Ser
 195 200 205

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Phe Arg
 210 215 220

Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ser Trp Asp Tyr Thr Leu
 225 230 235 240

Ser Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Ala
 245 250 255

Ala Ala His His His His His His Gly Gly Gly Gly Cys
 260 265

<210> 37

<211> 807

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human/murine
 chimeric single chain binding polypeptide (C6ML3-9
 sFv')

<400> 37

cagggtgcagc tgggtgcagtc tgggggcagag gtgaaaaagc ccgggggagtc tctgaagatc 60
 tctgtgaagg gttctggata cagctttacc agctactgga tcgcctgggt gcgccagatg 120

cccgggaaag gcctggagta catggggctc atctatcctg gtgactctga caccaaatac 180
 agcccgtcct tccaaggcca ggtcaccatc tcagtcgaca agtccgtcag cactgcctac 240
 ttgcaatgga gcagtctgaa gccctcggac agcgccgtgt atttttgtgc gagacatgac 300
 gtgggatatt gcagtagttc caactgcgca aagtggcctg aatacttcca gcattggggc 360
 cagggcaccc tggtcaccgt ctccctcaggt ggaggcgggt caggcggagg tggctctggc 420
 ggtggcggat cgcagtctgt gttgacgcag ccgccctcag tgtctgcggc cccaggacag 480
 aaggtcacca tctcctgctc tggaagcagc tccaacattg ggaataatta tgtatcctgg 540
 taccagcagc tccaggaac agcccccaaa ctccctcatct atgatcacac caatcggccc 600
 gcaggggtcc ctgaccgatt ctctggctcc aagtctggca cctcagcctc cctggccatc 660
 agtgggttcc ggtccgagga tgaggctgat tattactgtg cctcctggga ctacaccctc 720
 tcgggctggg tggtcggcgg aggaaccaag ctgaccgtcc taggtgcggc cgcacaccat 780
 catcaccatc acggtggtgg cggctgc 807

<210> 38

<211> 282

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human/murine
 chimeric single chain binding polypeptide
 (C6ML-3-9sFv'-L1-KDEL)

<400> 38

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
 1 5 10 15

Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe Thr Ser Tyr
 20 25 30

Trp Ile Ala Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Tyr Met
 35 40 45

Gly Leu Ile Tyr Pro Gly Asp Ser Asp Thr Lys Tyr Ser Pro Ser Phe
 50 55 60

Gln Gly Gln Val Thr Ile Ser Val Asp Lys Ser Val Ser Thr Ala Tyr
 65 70 75 80

B2
Cont

Leu Gln Trp Ser Ser Leu Lys Pro Ser Asp Ser Ala Val Tyr Phe Cys
 85 90 95

Ala Arg His Asp Val Gly Tyr Cys Ser Ser Ser Asn Cys Ala Lys Trp
 100 105 110

Pro Glu Tyr Phe Gln His Trp Gly Gln Gly Thr Leu Val Thr Val Ser
 115 120 125

Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
 130 135 140

Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Ala Ala Pro Gly Gln
 145 150 155 160

Lys Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Asn Asn
 165 170 175

Tyr Val Ser Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
 180 185 190

Ile Tyr Asp His Thr Asn Arg Pro Ala Gly Val Pro Asp Arg Phe Ser
 195 200 205

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Phe Arg
 210 215 220

Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ser Trp Asp Tyr Thr Leu
 225 230 235 240

Ser Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Ala
 245 250 255

Ala Ala His His His His His His Gly Gly Gly Gly Cys Leu Glu Ser
 260 265 270

Ser Ser Ser Gly Ser Glu Lys Asp Glu Leu

275

280

<210> 39

<211> 846

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Human/murine
chimeric single chain binding polypeptide
(C6ML-3-9sFv'-L1-KDEL)

<400> 39

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caggtgcagc tgggtgcagtc tgggggcagag gtgaaaaagc ccgggggagtc tctgaagatc 60
tcctgtaagg gttctggata cagctttacc agctactgga tcgcctgggt gcgccagatg 120
cccgggaaag gcctggagta catggggctc atctatcctg gtgactctga caccaaatac 180
agcccgtcct tccaaggcca ggtcaccatc tcagtcgaca agtccgtcag cactgcctac 240
ttgcaatgga gcagtctgaa gccctcggac agcgccgtgt atttttgtgc gagacatgac 300
gtgggatatt gcagtagttc caactgcgca aagtggcctg aatacttcca gcattggggc 360
cagggcaccc tggtcaccgt ctctcaggt ggaggcggtt caggcggagg tggctctggc 420
ggtggcggat cgcagtctgt gttgacgcag ccgccctcag tgtctgcggc cccaggacag 480
aaggtcacca tctcctgctc tggaagcagc tccaacattg ggaataatta tgtatcctgg 540
taccagcagc tcccaggaac agccccaaa ctctcatct atgatcacac caatcggcc 600
gcaggggtcc ctgaccgatt ctctggctcc aagtctggca cctcagcctc cctggccatc 660
agtgggttcc ggtccgagga tgaggctgat tattactgtg cctcctggga ctacaccctc 720
tcgggctggg tgttcggcgg aggaaccaag ctgaccgtcc taggtgcggc cgcacaccat 780
catcaccatc acggtggtgg cggctgcctc gagtcctcta gctctggatc cgaaaaagat 840
gaactg

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846

<210> 40

<211> 287

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Human/murine
chimeric single chain binding polypeptide
(C6ML3-9sFv'-L2-KDEL)

<400> 40

Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
 1 5 10 15

Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe Thr Ser Tyr
 20 25 30

Trp Ile Ala Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Tyr Met
 35 40 45

Gly Leu Ile Tyr Pro Gly Asp Ser Asp Thr Lys Tyr Ser Pro Ser Phe
 50 55 60

Gln Gly Gln Val Thr Ile Ser Val Asp Lys Ser Val Ser Thr Ala Tyr
 65 70 75 80

Leu Gln Trp Ser Ser Leu Lys Pro Ser Asp Ser Ala Val Tyr Phe Cys
 85 90 95

Ala Arg His Asp Val Gly Tyr Cys Ser Ser Ser Asn Cys Ala Lys Trp
 100 105 110

Pro Glu Tyr Phe Gln His Trp Gly Gln Gly Thr Leu Val Thr Val Ser
 115 120 125

Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
 130 135 140

Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Ala Ala Pro Gly Gln
 145 150 155 160

Lys Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Asn Asn
 165 170 175

Tyr Val Ser Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
 180 185 190

B2
Cont

Ile Tyr Asp His Thr Asn Arg Pro Ala Gly Val Pro Asp Arg Phe Ser
 195 200 205

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Phe Arg
 210 215 220

Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ser Trp Asp Tyr Thr Leu
 225 230 235 240

Ser Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Ala
 245 250 255

Ala Ala His His His His His His Gly Gly Gly Gly Cys Leu Glu Ser
 260 265 270

Ser Ser Ser Gly Ser Ser Ser Ser Gly Ser Glu Lys Asp Glu Leu
 275 280 285

<210> 41

<211> 861

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Human/murine
 chimeric single chain binding polypeptide
 (C6ML3-9sFv'-L2-KDEL)

<400> 41

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 cccgggaaag gcctggagta catggggctc atctatcctg gtgactctga caccaaatac 180
 agcccgctcct tccaaggcca ggtcaccatc tcagtcgaca agtccgtcag cactgcctac 240
 ttgcaatgga gcagtctgaa gccctcggac agcgccgtgt atttttgtgc gagacatgac 300
 gtgggatatt gcagtagttc caactgcgca aagtggcctg aatacttcca gcattggggc 360
 cagggcaccc tggtcaccgt ctctcaggt ggaggcgggt caggcggagg tggctctggc 420
 ggtggcggat cgcagtctgt gttgacgcag ccgcctcag tgtctgcggc cccaggacag 480

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aaggtcacca tctcctgctc tggaagcagc tccaacattg ggaataatta tgtatcctgg 540
taccagcagc tcccaggaac agcccccaaa ctctcatct atgatcacac caatcggccc 600
gcaggggtcc ctgaccgatt ctctggctcc aagtctggca cctcagcctc cctggccatc 660
agtgggttcc ggtccgagga tgaggctgat tattactgtg cctcctggga ctacaccctc 720
tcgggctggg tggtcggcgg aggaaccaag ctgaccgtcc taggtgcggc cgcacaccat 780
catcaccatc acggtggtgg cggctgcctc gagtctagca gctccggttc ctctagctct 840
ggatccgaaa aagatgaact g                                     861

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<210> 42

<211> 296

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Human/murine
chimeric single chain binding polypeptide
(C6ML3-9sFv'-L2-H14)

<400> 42

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Gln Val Gln Leu Val Gln Ser Gly Ala Glu Val Lys Lys Pro Gly Glu
  1              5              10              15

```

```

Ser Leu Lys Ile Ser Cys Lys Gly Ser Gly Tyr Ser Phe Thr Ser Tyr
      20              25              30

```

```

Trp Ile Ala Trp Val Arg Gln Met Pro Gly Lys Gly Leu Glu Tyr Met
    35              40              45

```

```

Gly Leu Ile Tyr Pro Gly Asp Ser Asp Thr Lys Tyr Ser Pro Ser Phe
    50              55              60

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```

Gln Gly Gln Val Thr Ile Ser Val Asp Lys Ser Val Ser Thr Ala Tyr
    65              70              75              80

```

```

Leu Gln Trp Ser Ser Leu Lys Pro Ser Asp Ser Ala Val Tyr Phe Cys
      85              90              95

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```

Ala Arg His Asp Val Gly Tyr Cys Ser Ser Ser Asn Cys Ala Lys Trp
    100              105              110

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Pro Glu Tyr Phe Gln His Trp Gly Gln Gly Thr Leu Val Thr Val Ser
115 120 125

Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser
130 135 140

Gln Ser Val Leu Thr Gln Pro Pro Ser Val Ser Ala Ala Pro Gly Gln
145 150 155 160

Lys Val Thr Ile Ser Cys Ser Gly Ser Ser Ser Asn Ile Gly Asn Asn
165 170 175

Tyr Val Ser Trp Tyr Gln Gln Leu Pro Gly Thr Ala Pro Lys Leu Leu
180 185 190

Ile Tyr Asp His Thr Asn Arg Pro Ala Gly Val Pro Asp Arg Phe Ser
195 200 205

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Phe Arg
210 215 220

Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ser Trp Asp Tyr Thr Leu
225 230 235 240

Ser Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Ala
245 250 255

Ala Ala His His His His His His Gly Gly Gly Gly Cys Leu Glu Ser
260 265 270

Ser Ser Ser Gly Ser Ser Ser Ser Gly Ser Lys Lys Ser Ala Lys Lys
275 280 285

Thr Pro Lys Lys Ala Lys Lys Pro
290 295

B2
Cont

<210> 43

<211> 888

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Human/murine
chimeric single chain binding polypeptide
(C6ML3-9sFv'-L2-H14)

<400> 43

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tcctgtaagg gttctggata cagctttacc agctactgga tcgcctgggt gcgccagatg 120
cccgggaaag gcctggagta catggggctc atctatcctg gtgactctga caccaaatac 180
agcccgctct tccaaggcca ggtcaccatc tcagtcgaca agtccgtcag cactgcctac 240
ttgcaatgga gcagtctgaa gccctcggac agcgccgtgt atttttgtgc gagacatgac 300
gtgggatatt gcagtagttc caactgcgca aagtggcctg aatacttcca gcattggggc 360
cagggcaccc tggtcaccgt ctctcaggt ggaggcggtt caggcggagg tggctctggc 420
gggtggcggat cgcagtctgt gttgacgcag ccgcctcag tgtctgcggc ccaggacag 480
aaggtcacca tctcctgctc tggaagcagc tccaacattg ggaataatta tgtatcctgg 540
taccagcagc tcccaggaac agcccccaa ctcctcatct atgatcacac caatcggccc 600
gcaggggtcc ctgaccgatt ctctggctcc aagtctggca cctcagcctc cctggccatc 660
agtgggttcc ggtccgagga tgaggctgat tattactgtg cctcctggga ctacaccctc 720
tcgggctggg tggtcggcgg aggaaccaag ctgaccgtcc taggtgcggc cgcacaccat 780
catcaccatc acggtggtgg cggctgcctc gagtctagca gctccggttc ctctagctct 840
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<210> 44

<211> 291

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Human/murine
chimeric single chain binding polypeptide
(C6ML3-9sFv'-L2-nls)

<400> 44

Gln	Val	Gln	Leu	Val	Gln	Ser	Gly	Ala	Glu	Val	Lys	Lys	Pro	Gly	Glu	1	5	10	15
Ser	Leu	Lys	Ile	Ser	Cys	Lys	Gly	Ser	Gly	Tyr	Ser	Phe	Thr	Ser	Tyr	20	25	30	
Trp	Ile	Ala	Trp	Val	Arg	Gln	Met	Pro	Gly	Lys	Gly	Leu	Glu	Tyr	Met	35	40	45	
Gly	Leu	Ile	Tyr	Pro	Gly	Asp	Ser	Asp	Thr	Lys	Tyr	Ser	Pro	Ser	Phe	50	55	60	
Gln	Gly	Gln	Val	Thr	Ile	Ser	Val	Asp	Lys	Ser	Val	Ser	Thr	Ala	Tyr	65	70	75	80
Leu	Gln	Trp	Ser	Ser	Leu	Lys	Pro	Ser	Asp	Ser	Ala	Val	Tyr	Phe	Cys	85	90	95	
Ala	Arg	His	Asp	Val	Gly	Tyr	Cys	Ser	Ser	Ser	Asn	Cys	Ala	Lys	Trp	100	105	110	
Pro	Glu	Tyr	Phe	Gln	His	Trp	Gly	Gln	Gly	Thr	Leu	Val	Thr	Val	Ser	115	120	125	
Ser	Gly	Gly	Gly	Gly	Ser	Gly	Gly	Gly	Gly	Ser	Gly	Gly	Gly	Gly	Ser	130	135	140	
Gln	Ser	Val	Leu	Thr	Gln	Pro	Pro	Ser	Val	Ser	Ala	Ala	Pro	Gly	Gln	145	150	155	160
Lys	Val	Thr	Ile	Ser	Cys	Ser	Gly	Ser	Ser	Ser	Asn	Ile	Gly	Asn	Asn	165	170	175	
Tyr	Val	Ser	Trp	Tyr	Gln	Gln	Leu	Pro	Gly	Thr	Ala	Pro	Lys	Leu	Leu	180	185	190	
Ile	Tyr	Asp	His	Thr	Asn	Arg	Pro	Ala	Gly	Val	Pro	Asp	Arg	Phe	Ser	195	200	205	

Gly Ser Lys Ser Gly Thr Ser Ala Ser Leu Ala Ile Ser Gly Phe Arg
 210 215 220

Ser Glu Asp Glu Ala Asp Tyr Tyr Cys Ala Ser Trp Asp Tyr Thr Leu
 225 230 235 240

Ser Gly Trp Val Phe Gly Gly Gly Thr Lys Leu Thr Val Leu Gly Ala
 245 250 255

Ala Ala His His His His His His Gly Gly Gly Gly Cys Leu Glu Ser
 260 265 270

Ser Ser Ser Gly Ser Ser Ser Ser Gly Ser Thr Pro Pro Lys Lys Lys
 275 280 285

Arg Lys Val
 290

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<211> 873

<212> DNA

<213> Artificial Sequence

<220>

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 (C6ML3-9sFv'-L2-nls)

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 cagggcaccc tggtcaccgt ctccctcaggt ggaggcgggt caggcggagg tggctctggc 420
 ggtggcggat cgcagtctgt gttgacgcag ccgcctcag tgtctgcggc ccaggacag 480


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<210> 47
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Gly Gly Gly Gly Ser
1 5

B2
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